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## What is claimed is:

- A method for the detection of one or more NF-κB regulatory factors comprising the steps of:
  - a) providing a slimb protein, and a sample suspected of containing one or more NF-kB regulatory factors; and
  - exposing said slimb protein to said sample under conditions such that said slimb protein binds to said one or more NF-κB regulatory factors to form a slimb/regulatory factor complex.
- The method of Claim 1, further comprising the step of detecting said slimb/regulatory factor complex.
- The method of Claim 1, further comprising the step of observing said slimb/regulatory factor complex for degradation of said one or more NF-κB regulatory factors.
- 4. The method of Claim 1, further comprising the step of exposing said slimb protein and one or more NF-κB regulatory factors to an F-box protein antagonist.
- The method of Claim 4, wherein said F-box protein antagonist prevents the formation of said slimb/regulatory factor complex.

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- A method for the detection of a slimb protein complex, comprising the steps of:
  - a) providing a slimb protein and a sample suspected of containing one or more proteins capable of forming a complex with said slimb protein;
    and
  - b) exposing said slimb protein to said one or more proteins capable of forming a complex with said slimb protein under conditions such that said slimb protein binds to said one or more proteins capable of forming a complex with said slimb protein to form a slimb protein complex.
- The method of Claim 6, further comprising the step of detecting said slimb protein complex.
- 8. The method of Claim 6, wherein step b) further comprises exposing said slimb protein and said one or more proteins capable of forming a complex with said slimb protein to an F-box protein antagonist.
- The method of Claim 8, wherein said F-box protein antagonist prevents the formation of said slimb protein complex.
- 10. An isolated nucleotide sequence comprising nucleotide sequence encoding at least one functionally active fragement of an F-box protein, wherein said sequence consists of a least a portion of the sequence set forth in SEQ ID NOS: 54 and 56.

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